

## SEQUENCE LISTING

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<110> Hedley, Mary Lynne
      Urban, Robert G.
      Chicz, Roman M.
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<213> Human Papilloma virus
<400> 152
Arg Arg Pro Tyr Lys Leu Pro Asp Leu Cys Thr Glu Leu Asn Thr Ser
                                     10
Leu Gln Asp Ile Glu Ile Thr Cys Val Tyr Cys Lys Thr Val Leu Glu
                                                      30
                                 25
             20
Leu Thr Glu Val Phe Glu Phe Ala Phe Lys
         35
<210> 153
<211> 27
<212> PRT
<213> Human Papilloma virus
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<400> 153
Ser Val Tyr Gly Asp Thr Leu Glu Lys Leu Thr Asn Thr Gly Leu Tyr
                                     10
Asn Leu Leu Ile Arg Cys Leu Arg Cys Gln Lys
<210> 154
<211> 19
<212> PRT
<213> Human Papilloma virus
<400> 154
Lys Ala Thr Leu Gln Asp Ile Val Leu His Leu Glu Pro Gln Asn Glu
Ile Pro Val
<210> 155
<211> 14
<212> PRT
<213> Human Papilloma virus
<400> 155
His Thr Met Leu Cys Met Cys Cys Lys Cys Glu Ala Arg Ile
<210> 156
<211> 17
<212> PRT
<213> Human Papilloma virus
<400> 156
Ala Phe Gln Gln Leu Phe Leu Asn Thr Leu Ser Phe Val Cys Pro Trp
                  5
Cys
<210> 157
<211> 236
<212> PRT
<213> Artificial Sequence
<220>
<223> Artificial fusion sequence
<400> 157
Ala Met Phe Gln Asp Pro Gln Glu Arg Pro Arg Lys Leu Pro Gln Leu
                                     10
Cys Thr Glu Leu Leu Arg Arg Glu Val Tyr Asp Phe Ala Phe Arg
                                 25
                                                      30
             20
Asp Leu Cys Ile Val Tyr Arg Asp Gly Asn Pro Tyr Lys Ile Ser Glu
                                                 45
         35
                             40
Tyr Arg His Tyr Cys Tyr Ser Leu Tyr Gly Thr Thr Leu Glu Gln Gln
    5Ō
                         55
Tyr Asn Lys Thr Leu His Glu Tyr Met Leu Asp Leu Gln Pro Glu Thr
                     70
                                         75
Thr Asp Leu Tyr Ser Tyr Gln Ala Glu Pro Asp Arg Ala His Tyr Asn
                                     90
                85
Ile Val Thr Phe Leu Leu Met Gly Thr Leu Gly Ile Val Cys Pro Ile
                                105
                                                     110
            100
Cys Ser Gln Lys Pro Arg Arg Pro Tyr Lys Leu Pro Asp Leu Cys Thr
                            120
        115
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Glu Leu Asn Thr Ser Leu Gln Asp Ile Glu Ile Thr Cys Val Tyr Cys
                                            140
                        135
   130
Lys Thr Val Leu Glu Leu Thr Glu Val Phe Glu Phe Ala Phe Lys Ser
                    150
                                       155
Val Tyr Gly Asp Thr Leu Glu Lys Leu Thr Asn Thr Gly Leu Tyr Asn
                                  170
               165
Leu Leu Ile Arg Cys Leu Arg Cys Gln Lys Lys Ala Thr Leu Gln Asp
                              185
           180
Ile Val Leu His Leu Glu Pro Gln Asn Glu Ile Pro Val His Thr Met
                           200
                                              205
       195
Leu Cys Met Cys Cys Lys Cys Glu Ala Arg Ile Ala Phe Gln Gln Leu
                       215
                                           220
Phe Leu Asn Thr Leu Ser Phe Val Cys Pro Trp Cys
                   230
<210> 158
<211> 237
<212> PRT
<213> Artificial Sequence
<223> Artificial fusion sequence
<400> 158
Met Ala Met Phe Gln Asp Pro Gln Glu Arg Pro Arg Lys Leu Pro Gln
                                    10
Leu Cys Thr Glu Leu Leu Arg Arg Glu Val Tyr Asp Phe Ala Phe
           20
                                25
Arg Asp Leu Cys Ile Val Tyr Arg Asp Gly Asn Pro Tyr Lys Ile Ser
        35
                            40
Glu Tyr Arg His Tyr Cys Tyr Ser Leu Tyr Gly Thr Thr Leu Glu Gln
    50
                        55
Gln Tyr Asn Lys Thr Leu His Glu Tyr Met Leu Asp Leu Gln Pro Glu
                    70
                                        75
Thr Thr Asp Leu Tyr Ser Tyr Gln Ala Glu Pro Asp Arg Ala His Tyr
                85
                                    90
Asn Ile Val Thr Phe Leu Leu Met Gly Thr Leu Gly Ile Val Cys Pro
                               105
                                                   110
           100
Ile Cys Ser Gln Lys Pro Arg Arg Pro Tyr Lys Leu Pro Asp Leu Cys
                           120
                                                125
       115
Thr Glu Leu Asn Thr Ser Leu Gln Asp Ile Glu Ile Thr Cys Val Tyr
                       135
                                           140
   130
Cys Lys Thr Val Leu Glu Leu Thr Glu Val Phe Glu Phe Ala Phe Lys
                                       155
                  150
Ser Val Tyr Gly Asp Thr Leu Glu Lys Leu Thr Asn Thr Gly Leu Tyr
                                  170
               165
Asn Leu Leu Ile Arg Cys Leu Arg Cys Gln Lys Lys Ala Thr Leu Gln
           180
                               185
Asp Ile Val Leu His Leu Glu Pro Gln Asn Glu Ile Pro Val His Thr
                           200
                                               205
Met Leu Cys Met Cys Cys Lys Cys Glu Ala Arg Ile Ala Phe Gln Gln
                      215
                                           220
Leu Phe Leu Asn Thr Leu Ser Phe Val Cys Pro Trp Cys
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<210> 159
<211> 119
<212> PRT
<213> Artificial Sequence
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<223> Artificial fusion sequence

<400> 159 Arg Arg Pro Tyr Lys Leu Pro Asp Leu Cys Thr Glu Leu Asn Thr Ser Leu Gln Asp Ile Glu Ile Thr Cys Val Tyr Cys Lys Thr Val Leu Glu 25 Leu Thr Glu Val Phe Glu Phe Ala Phe Lys Ser Val Tyr Gly Asp Thr 40 35 Leu Glu Lys Leu Thr Asn Thr Gly Leu Tyr Asn Leu Leu Ile Arg Cys 55 Leu Arg Cys Gln Lys Lys Ala Thr Leu Gln Asp Ile Val Leu His Leu 70 75 Glu Pro Gln Asn Glu Ile Pro Val His Thr Met Leu Cys Met Cys Cys 90 85 Lys Cys Glu Ala Arg Ile Ala Phe Gln Gln Leu Phe Leu Asn Thr Leu 100 105 Ser Phe Val Cys Pro Trp Cys 115 <210> 160 <211> 261 <212> PRT <213> Artificial Sequence <220> <223> Artificial fusion sequence <400> 160 Met Ala Ile Ser Gly Val Pro Val Leu Gly Phe Phe Ile Ile Ala Val 10 Leu Met Ser Ala Gln Glu Ser Trp Ala Ala Met Phe Gln Asp Pro Gln 25 Glu Arg Pro Arg Lys Leu Pro Gln Leu Cys Thr Glu Leu Leu Arg 35 40 45 Arg Glu Val Tyr Asp Phe Ala Phe Arg Asp Leu Cys Ile Val Tyr Arg 55 Asp Gly Asn Pro Tyr Lys Ile Ser Glu Tyr Arg His Tyr Cys Tyr Ser 70 Leu Tyr Gly Thr Thr Leu Glu Gln Gln Tyr Asn Lys Thr Leu His Glu 90 85 Tyr Met Leu Asp Leu Gln Pro Glu Thr Thr Asp Leu Tyr Ser Tyr Gln 105 Ala Glu Pro Asp Arg Ala His Tyr Asn Ile Val Thr Phe Leu Leu Met 115 120 125 Gly Thr Leu Gly Ile Val Cys Pro Ile Cys Ser Gln Lys Pro Arg Arg 135 140 Pro Tyr Lys Leu Pro Asp Leu Cys Thr Glu Leu Asn Thr Ser Leu Gln 150 155 Asp Ile Glu Ile Thr Cys Val Tyr Cys Lys Thr Val Leu Glu Leu Thr 170 165 Glu Val Phe Glu Phe Ala Phe Lys Ser Val Tyr Gly Asp Thr Leu Glu 180 185 190 Lys Leu Thr Asn Thr Gly Leu Tyr Asn Leu Leu Ile Arg Cys Leu Arg 195 200 205 Cys Gln Lys Lys Ala Thr Leu Gln Asp Ile Val Leu His Leu Glu Pro

215

230

Gln Asn Glu Ile Pro Val His Thr Met Leu Cys Met Cys Cys Lys Cys

Glu Ala Arg Ile Ala Phe Gln Gln Leu Phe Leu Asn Thr Leu Ser Phe

220

235

245 Val Cys Pro Trp Cys 260

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<210> 161
<211> 340
<212> DNA
<213> Artificial Sequence
<223> Artificial fusion sequence
<400> 161
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                                                                           60
ctectectge geogegaggt gtatgaette gettttegeg acetgtgeat egtetacege
                                                                          120
                                                                          180
gatggcaacc catataagat ctccgaqtac cgccactact gctattccct gtatggcacc
                                                                          240
accetqqaqe aqcaqtacaa caagaceetg caegagtaca tgetggatet geageeagag
accaccgate tgtactecta teaggetgag ceagategeg etcactataa categtgace
                                                                          300
ttcctgctga tgggcaccct gggcatcgtg tgcccaatct
                                                                          340
<210> 162
<211> 786
<212> DNA
<213> Artificial Sequence
<220>
<223> Artificial fusion sequence
atggetatet ceggegtgee agtgetggge ttetteatea tegetgtget gatgteeget
                                                                           60
caggagtect gggetgetat gttecaggac ecacaggaac geccaegcaa actgecacag
                                                                          120
ctgtgcaccg agetectect gegeegegag gtgtatgact tegettiteg egacetgtge
                                                                          180
atogtotaco gogatggoaa cocatataag atotoogagt acogcoacta otgotattoo ctgtatggoa coaccotgga goagcagtac aacaagacco tgcacgagta catgotggat
                                                                          240
                                                                          300
ctgcagccag agaccaccga totgtactcc tatcaggctg agccagatcg cgctcactat
                                                                          360
aacatcgtga cettectget gatgggcace etgggcatcg tgtgcccaat etgeteccag
                                                                          420
                                                                          480
aageecegge gaccetacaa getaectgat etgtgcaegg aactgaacae tteactgcaa
                                                                          540
gacatagaaa taacctgtgt atattgcaag acagtattgg aacttacaga ggtatttgaa
tttgcattta aatctgtgta tggagacaca ttggaaaaac taactaacac tgggttatac
                                                                          600
aatitattaa taaggigcct gcggtgccag aaaaaggcaa cattgcaaga cattgtattg
                                                                          660
catttagagc cccaaaatga aattccggtt cacacaatgt tgtgtatgtg ttgtaagtgt
                                                                          720
                                                                          780
qaaqccaqaa ttqcattcca gcaqctqttt ctgaacaccc tqtcctttqt gtqtccgtgg
                                                                          786
tgttaa
<210> 163
<211> 9
<212> PRT
<213> Human Papilloma virus
<400> 163
Glu Val Tyr Asp Phe Ala Phe Arg Asp
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